

<110> Oregon Health & Science University Michael, Heinrich Charles Corless, Christopher Lee Fletcher, Jonathan Alfred Demetri, George D. <120> ACTIVATING MUTATIONS OF PLATELET DERIVED GROWTH FACTOR RECEPTOR ALPHA (PDGFRA) AS DIAGNOSTIC MARKERS AND THERAPEUTIC TARGETS <130> 899-65892-02 <150> US 60/389,107 <151> 2002-06-13 <150> US 60/438,899 <151> 2003-01-08 <150> PCT/US2003/18907 <151> 2003-06-13 <160> 27 <170> PatentIn version 3.2 <210> 1 <211> 6633 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (395)..(3664) <400> 1 ttctccccgc cccccagttg ttgtcgaagt ctgggggttg ggactggacc ccctgattgc 60 gtaagagcaa aaagcgaagg cgcaatctgg acactgggag attcggagcg cagggagttt 120 gagagaaact tttattttga agagaccaag gttgaggggg ggcttatttc ctgacagcta 180 tttacttaga gcaaatgatt agttttagaa ggatggacta taacattgaa tcaattacaa 240 aacgcggttt ttgagcccat tactgttgga gctacaggga gagaaacagg aggagactgc 300 aagagatcat ttgggaaggc cgtgggcacg ctctttactc catgtgtggg acattcattg 360

ttc ctg gtc tta ggc tgt ctt ctc aca ggg ctg agc cta atc ctc tgc 463 Phe Leu Val Leu Gly Cys Leu Leu Thr Gly Leu Ser Leu Ile Leu Cys 10

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415

511

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Cys Phe Gly Glu Ser Glu Val Ser Trp Gln. Tyr Pro Met Ser Glu Glu 50 60

Glu Ser Ser Asp Val Glu Ile Arg Asn Glu Glu Asn Asn Ser Gly Leu 65 70 75 80

Phe Val Thr Val Leu Glu Val Ser Ser Ala Ser Ala Ala His Thr Gly 85 90 95

Leu Tyr Thr Cys Tyr Tyr Asn His Thr Gln Thr Glu Glu Asn Glu Leu 100 105 110

Glu Gly Arg His Ile Tyr Ile Tyr Val Pro Asp Pro Asp Val Ala Phe
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Val Pro Leu Gly Met Thr Asp Tyr Leu Val Ile Val Glu Asp Asp Asp 130 135 140

Ser Ala Ile Ile Pro Cys Arg Thr Thr Asp Pro Glu Thr Pro Val Thr 145 150 155 160

Leu His Asn Ser Glu Gly Val Val Pro Ala Ser Tyr Asp Ser Arg Gln
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Gly Phe Asn Gly Thr Phe Thr Val Gly Pro Tyr Ile Cys Glu Ala Thr 180 185 190

Val Lys Gly Lys Lys Phe Gln Thr Ile Pro Phe Asn Val Tyr Ala Leu Lys Ala Thr Ser Glu Leu Asp Leu Glu Met Glu Ala Leu Lys Thr Val Tyr Lys Ser Gly Glu Thr Ile Val Val Thr Cys Ala Val Phe Asn Asn Glu Val Val Asp Leu Gln Trp Thr Tyr Pro Gly Glu Val Lys Gly Lys Gly Ile Thr Met Leu Glu Glu Ile Lys Val Pro Ser Ile Lys Leu Val Tyr Thr Leu Thr Val Pro Glu Ala Thr Val Lys Asp Ser Gly Asp Tyr Glu Cys Ala Ala Arg Gln Ala Thr Arg Glu Val Lys Glu Met Lys Lys Val Thr Ile Ser Val His Glu Lys Gly Phe Ile Glu Ile Lys Pro Thr Phe Ser Gln Leu Glu Ala Val Asn Leu His Glu Val Lys His Phe Val Val Glu Val Arg Ala Tyr Pro Pro Pro Arg Ile Ser Trp Leu Lys Asn Asn Leu Thr Leu Ile Glu Asn Leu Thr Glu Ile Thr Thr Asp Val Glu Lys Ile Gln Glu Ile Arg Tyr Arg Ser Lys Leu Lys Leu Ile Arg Ala Lys Glu Glu Asp Ser Gly His Tyr Thr Ile Val Ala Gln Asn Glu Asp Ala Val Lys Ser Tyr Thr Phe Glu Leu Leu Thr Gln Val Pro Ser Ser Ile Leu Asp Leu Val Asp Asp His His Gly Ser Thr Gly Gly Gln Thr

Val Arg Cys Thr Ala Glu Gly Thr Pro Leu Pro Asp Ile Glu Trp Met Ile Cys Lys Asp Ile Lys Lys Cys Asn Asn Glu Thr Ser Trp Thr Ile Leu Ala Asn Asn Val Ser Asn Ile Ile Thr Glu Ile His Ser Arg Asp Arg Ser Thr Val Glu Gly Arg Val Thr Phe Ala Lys Val Glu Glu Thr Ile Ala Val Arg Cys Leu Ala Lys Asn Leu Leu Gly Ala Glu Asn Arg Glu Leu Lys Leu Val Ala Pro Thr Leu Arg Ser Glu Leu Thr Val Ala Ala Ala Val Leu Val Leu Val Ile Val Ile Ser Leu Ile Val Leu Val Val Ile Trp Lys Gln Lys Pro Arg Tyr Glu Ile Arg Trp Arg Val Ile Glu Ser Ile Ser Pro Asp Gly His Glu Tyr Ile Tyr Val Asp Pro Met Gln Leu Pro Tyr Asp Ser Arg Trp Glu Phe Pro Arg Asp Gly Leu Val Leu Gly Arg Val Leu Gly Ser Gly Ala Phe Gly Lys Val Val Glu Gly Thr Ala Tyr Gly Leu Ser Arg Ser Gln Pro Val Met Lys Val Ala Val Lys Met Leu Lys Pro Thr Ala Arg Ser Ser Glu Lys Gln Ala Leu Met Ser Glu Leu Lys Ile Met Thr His Leu Gly Pro His Leu Asn Ile Val Asn Leu Leu Gly Ala Cys Thr Lys Ser Gly Pro Ile Tyr Ile

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Ala	Asp	Thr	Thr 740	Gln	Tyr	Val	Pro	Met 745	Leu	Glu	Arg	Lys	Glu 750	Val	Ser
Lys	Tyr	Ser 755	Asp	Ile	Gln	Arg	Ser 760	Leu	Tyr	Asp	Arg	Pro 765	Ala	Ser	Tyr
Lys	Lys 770	Lys	Ser	Met	Leu	Asp 775	Ser	Glu	Val	Lys	Asn 780	Leu	Leu	Ser	Asp
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Lys	Ile	Cys 835	Asp	Phe	Gly	Leu	Ala 840	Arg	Asp	Ile	Met	His 845	Asp	Ser	Asn
Tyr	Val 850	Ser	Lys	Gly	Ser	Thr 855	Phe	Leu	Pro	Val	Lys 860	Trp	Met	Ala	Pro
Glu 865	Ser	Ile	Phe	Asp	Asn 870	Leu	Tyr	Thr	Thr	Leu 875	Ser	Asp	Val	Trp	Ser 880
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Pro	Gly	Met	Met 900	Val	Asp	Ser	Thr	Phe 905	Tyr	Asn	Lys	Ile	Lys 910	Ser	Gly

Tyr Arg Met Ala Lys Pro Asp His Ala Thr Ser Glu Val Tyr Glu Ile 915 920 925

Met Val Lys Cys Trp Asn Ser Glu Pro Glu Lys Arg Pro Ser Phe Tyr 930 935 940

His Leu Ser Glu Ile Val Glu Asn Leu Leu Pro Gly Gln Tyr Lys Lys 945 950 955 960

Ser Tyr Glu Lys Ile His Leu Asp Phe Leu Lys Ser Asp His Pro Ala 965 970 975

Val Ala Arg Met Arg Val Asp Ser Asp Asn Ala Tyr Ile Gly Val Thr 980 985 990

Tyr Lys Asn Glu Glu Asp Lys Leu Lys Asp Trp Glu Gly Gly Leu Asp 995 1000 1005

Glu Gln Arg Leu Ser Ala Asp Ser Gly Tyr Ile Ile Pro Leu Pro 1010 1015 1020

Asp Ile Asp Pro Val Pro Glu Glu Glu Asp Leu Gly Lys Arg Asn 1025 1030 1035

Arg His Ser Ser Gln Thr Ser Glu Glu Ser Ala Ile Glu Thr Gly 1040 1050

Ser Ser Ser Ser Thr Phe Ile Lys Arg Glu Asp Glu Thr Ile Glu 1055 1060 1065

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Cys Phe Gly Glu Ser Glu Val Ser Trp Gln Tyr Pro Met Ser Glu Glu 50 55 60

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	cat His 425						_	_			_		_	_		1711
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Cys Phe Gly Glu Ser Glu Val Ser Trp Gln Tyr Pro Met Ser Glu Glu 50 55 60

Glu Ser Ser Asp Val Glu Ile Arg Asn Glu Glu Asn Asn Ser Gly Leu 65 70 75 80

Phe Val Thr Val Leu Glu Val Ser Ser Ala Ser Ala Ala His Thr Gly
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Leu Tyr Thr Cys Tyr Tyr Asn His Thr Gln Thr Glu Glu Asn Glu Leu 100 105 110

Glu Gly Arg His Ile Tyr Ile Tyr Val Pro Asp Pro Asp Val Ala Phe 115 120 125

Val Pro Leu Gly Met Thr Asp Tyr Leu Val Ile Val Glu Asp Asp Asp 130 135 140

Ser Ala Ile Ile Pro Cys Arg Thr Thr Asp Pro Glu Thr Pro Val Thr 145 150 155 160

Leu His Asn Ser Glu Gly Val Val Pro Ala Ser Tyr Asp Ser Arg Gln
165 170 175

Gly Phe Asn Gly Thr Phe Thr Val Gly Pro Tyr Ile Cys Glu Ala Thr 180 185 190

Val Lys Gly Lys Lys Phe Gln Thr Ile Pro Phe Asn Val Tyr Ala Leu 195 200 205

Lys Ala Thr Ser Glu Leu Asp Leu Glu Met Glu Ala Leu Lys Thr Val 210 215 220

Tyr Lys Ser Gly Glu Thr Ile Val Val Thr Cys Ala Val Phe Asn Asn 225 230 235 240

Glu Val Val Asp Leu Gln Trp Thr Tyr Pro Gly Glu Val Lys Gly Lys

Gly Ile Thr Met Leu Glu Glu Ile Lys Val Pro Ser Ile Lys Leu Val Tyr Thr Leu Thr Val Pro Glu Ala Thr Val Lys Asp Ser Gly Asp Tyr Glu Cys Ala Ala Arg Gln Ala Thr Arg Glu Val Lys Glu Met Lys Lys Val Thr Ile Ser Val His Glu Lys Gly Phe Ile Glu Ile Lys Pro Thr

Phe Ser Gln Leu Glu Ala Val Asn Leu His Glu Val Lys His Phe Val

Val Glu Val Arg Ala Tyr Pro Pro Pro Arg Ile Ser Trp Leu Lys Asn

Asn Leu Thr Leu Ile Glu Asn Leu Thr Glu Ile Thr Thr Asp Val Glu

Lys Ile Gln Glu Ile Arg Tyr Arg Ser Lys Leu Lys Leu Ile Arg Ala

Lys Glu Glu Asp Ser Gly His Tyr Thr Ile Val Ala Gln Asn Glu Asp

Ala Val Lys Ser Tyr Thr Phe Glu Leu Leu Thr Gln Val Pro Ser Ser

Ile Leu Asp Leu Val Asp Asp His His Gly Ser Thr Gly Gly Gln Thr

Val Arg Cys Thr Ala Glu Gly Thr Pro Leu Pro Asp Ile Glu Trp Met

Ile Cys Lys Asp Ile Lys Lys Cys Asn Asn Glu Thr Ser Trp Thr Ile

Leu Ala Asn Asn Val Ser Asn Ile Ile Thr Glu Ile His Ser Arq Asp

Arg Ser Thr Val Glu Gly Arg Val Thr Phe Ala Lys Val Glu Glu Thr

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Glu	Leu	Lys 515	Leu	Val	Ala	Pro	Thr 520	Leu	Arg	Ser	Glu	Leu 525	Thr	Val	Ala
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Pro	Met	Gln	Leu 580	Pro	Tyr	Asp	Ser	Arg 585	Trp	Glu	Phe	Pro	Arg 590	Asp	Gly
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Glu	Gly 610	Thr	Ala	Tyr	Gly	Leu 615	Ser	Arg	Ser	Gln	Pro 620	Val	Met	Lys	Val
Ala 625	Val	Lys	Met	Leu	Lys 630	Pro	Thr	Ala	Arg	Ser 635	Ser	Glu	Lys	Gln	Ala 640
Leu	Met	Ser	Glu	Leu 645	Lys	Ile	Met	Thr	His 650	Leu	Gly	Pro	His	Leu 655	Asn
Ile	Val	Asn	Leu 660	Leu	Gly	Ala	Cys	Thr 665	Lys	Ser	Gly	Pro	Ile 670	Tyr	Ile
Ile	Thr	Glu 675	Tyr	Cys	Phe	Tyr	Gly 680	Asp	Leu	Val	Asn	Tyr 685	Leu	His	Lys

Val Ile Leu Ser Phe Glu Asn Asn Gly Asp Tyr Met Asp Met Lys Gln

Asn Arg Asp Ser Phe Leu Ser His His Pro Glu Lys Pro Lys Lys Glu

Leu Asp Ile Phe Gly Leu Asn Pro Ala Asp Glu Ser Thr Arg Ser Tyr

Glu Val Ser

				, 23								
Ala	Asp	Thr	Thr 740	Gln	Tyr	Val	Pro	Met 745	Leu	Glu	Arg	Lys

- Lys Tyr Ser Asp Ile Gln Arg Ser Leu Tyr Asp Arg Pro Ala Ser Tyr 755 760 765
- Lys Lys Ser Met Leu Asp Ser Glu Val Lys Asn Leu Leu Ser Asp 770 780
- Asp Asn Ser Glu Gly Leu Thr Leu Leu Asp Leu Leu Ser Phe Thr Tyr 785 790 795 800
- Gln Val Ala Arg Gly Met Glu Phe Leu Ala Ser Lys Asn Cys Val His 805 810 815
- Arg Asp Leu Ala Ala Arg Asn Val Leu Leu Ala Gln Gly Lys Ile Val 820 825 830
- Lys Ile Cys Asp Phe Gly Leu Ala Arg Asp Ser Asn Tyr Val Ser Lys 835 840 845
- Gly Ser Thr Phe Leu Pro Val Lys Trp Met Ala Pro Glu Ser Ile Phe 850 855 860
- Asp Asn Leu Tyr Thr Thr Leu Ser Asp Val Trp Ser Tyr Gly Ile Leu 865 870 875 880
- Leu Trp Glu Ile Phe Ser Leu Gly Gly Thr Pro Tyr Pro Gly Met Met 885 . 890 895
- Val Asp Ser Thr Phe Tyr Asn Lys Ile Lys Ser Gly Tyr Arg Met Ala 900 905 910
- Lys Pro Asp His Ala Thr Ser Glu Val Tyr Glu Ile Met Val Lys Cys 915 920 925
- Trp Asn Ser Glu Pro Glu Lys Arg Pro Ser Phe Tyr His Leu Ser Glu 930 935 940
- Ile Val Glu Asn Leu Leu Pro Gly Gln Tyr Lys Lys Ser Tyr Glu Lys 945 950 955 960
- Ile His Leu Asp Phe Leu Lys Ser Asp His Pro Ala Val Ala Arg Met

Arg Val Asp Ser Asp Asn Ala Tyr Ile Gly Val Thr Tyr Lys Asn Glu 980 985 990
Glu Asp Lys Leu Lys Asp Trp Glu Gly Gly Leu Asp Glu Gln Arg Leu 995 1000 1005
Ser Ala Asp Ser Gly Tyr Ile Ile Pro Leu Pro Asp Ile Asp Pro 1010 1015 1020
Val Pro Glu Glu Glu Asp Leu Gly Lys Arg Asn Arg His Ser Ser 1025 1030 1035
Gln Thr Ser Glu Glu Ser Ala Ile Glu Thr Gly Ser Ser Ser 1040 1045 1050
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cct gct gat gaa agc aca cgg agc tat gtt att tta tct ttt gaa aac

Pro Ala Asp Glu Ser Thr Arg Ser Tyr Val Ile Leu Ser Phe Glu Asn

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	_	gtc Val						_	_			_				2767
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		gac Asp														2959
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		gat Asp														3055
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		caa Gln			_	_		_				_	_		_	3295

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Cys Phe Gly Glu Ser Glu Val Ser Trp Gln Tyr Pro Met Ser Glu Glu 50 55 60

Glu Ser Ser Asp Val Glu Ile Arg Asn Glu Glu Asn Asn Ser Gly Leu 65 70 75 80

Phe Val Thr Val Leu Glu Val Ser Ser Ala Ser Ala Ala His Thr Gly 85 90 95

Leu Tyr Thr Cys Tyr Tyr Asn His Thr Gln Thr Glu Glu Asn Glu Leu 100 105 110

Glu Gly Arg His Ile Tyr Ile Tyr Val Pro Asp Pro Asp Val Ala Phe 115 120 125

Val Pro Leu Gly Met Thr Asp Tyr Leu Val Ile Val Glu Asp Asp Asp 130 135 140

Ser Ala Ile Ile Pro Cys Arg Thr Thr Asp Pro Glu Thr Pro Val Thr 145 150 155 160

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Val	ГАż	Gly 195	Lys	Lys	Phe	Gln	Thr 200	Ile	Pro	Phe	Asn	Val 205	Tyr	Ala	Leu
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Pro Met Gln	Leu Pro 580	Tyr Asp	Ser	Arg 585	Trp	Glu	Phe	Pro	Arg 590	Asp	Gly
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Glu Gly Thr 610	Ala Tyr	Gly Leu 615	Ser	Arg	Ser	Gln	Pro 620	Val	Met	Lys	Val
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Ile V	/al	Asn	Leu 660	Leu	Gly	Ala	Cys	Thr 665	Lys	Ser	Gly	Pro	Ile 670	Tyr	Ile
Ile T	Thr	Glu 675	Tyr	Cys	Phe	Tyr	Gly 680	Asp	Leu	Val	Asn	Tyr 685	Leu	His	Lys
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Lys T	Tyr	Ser 755	Asp	Ile	Gln	Arg	Ser 760	Leu	Tyr	Asp	Arg	Pro 765	Ala	Ser	Tyr
Lys I	Lys 770	Lys	Ser	Met	Leu	Asp 775	Ser	Glu	Val	Lys	Asn 780	Leu	Leu	Ser	Asp
Asp #	Asn	Ser	Glu	Gly	Leu 790	Thr	Leu	Leu	Asp	Leu 795	Leu	Ser	Phe	Thr	Tyr 800
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Lys 1	[le	Cys 835	Asp	Phe	Gly	Leu	Ala 840	Arg	Asp	Ile	Met	Pro 845	Tyr	Val	Ser
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Lys Ile His Leu Asp Phe Leu Lys Ser Asp His Pro Ala Val Ala Arg 965 970 975

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Glu Glu Asp Lys Leu Lys Asp Trp Glu Gly Gly Leu Asp Glu Gln Arg 995 1000 1005

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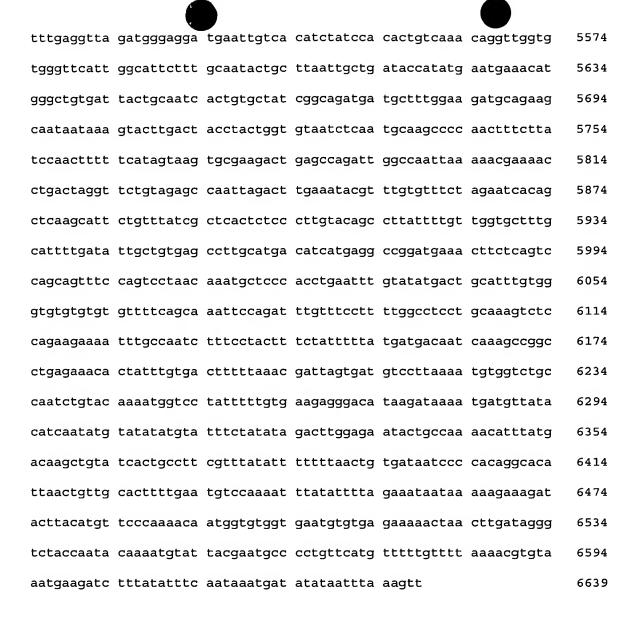
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Glu Cys Ala Ala Arg Gln Ala Thr Arg Glu Val Lys Glu Met Lys Lys

Tyr Thr Leu Thr Val Pro Glu Ala Thr Val Lys Asp Ser Gly Asp Tyr



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Phe Ser Gln Leu Glu Ala Val Asn Leu His Glu Val Lys His Phe Val 325 330 335

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Asn Leu Thr Leu Ile Glu Asn Leu Thr Glu Ile Thr Thr Asp Val Glu
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Lys Ile Gln Glu Ile Arg Tyr Arg Ser Lys Leu Lys Leu Ile Arg Ala 370 375 380

Lys Glu Glu Asp Ser Gly His Tyr Thr Ile Val Ala Gln Asn Glu Asp 385 390 395 400

Ala Val Lys Ser Tyr Thr Phe Glu Leu Leu Thr Gln Val Pro Ser Ser 405 410 415

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Ile Cys Lys Asp Ile Lys Lys Cys Asn Asn Glu Thr Ser Trp Thr Ile 450 455 460

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Glu Leu Lys Leu Val Ala Pro Thr Leu Arg Ser Glu Leu Thr Val Ala 515 520 525

Ala Ala Val Leu Val Leu Val Ile Val Ile Ser Leu Ile Val

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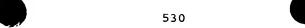
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109

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Thr Gly Gly Gln Thr Val Arg Cys Thr Ala Glu Gly Thr Pro Leu Pro 430 435

168408 gat att gag tgg atg ata tgc aaa gat att aag aa gtatggaaaa Asp Ile Glu Trp Met Ile Cys Lys Asp Ile Lys Lys 445

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gaa act tcc tgg act att ttg gcc aac aat gtc tca aac atc atc acg 169472 Glu Thr Ser Trp Thr Ile Leu Ala Asn Asn Val Ser Asn Ile Ile Thr 460 465 470	?
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ctt gga gct gag aac cga gag ctg aag ctg gtg gct ccc a gtgagttcct 169618 Leu Gly Ala Glu Asn Arg Glu Leu Lys Leu Val Ala Pro 510 515	}
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710 71

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Tyr Val Ile Leu Ser

	gaa Glu			_		_	_	_	_	_	_	_		176257
_	tat Tyr	_	_		_				_				_	176305

atc cag aga tca ctc tat gat cgt cca gcc tca tat aag aag aaa tct 176353

Ile Gln Arg Ser Leu Tyr Asp Arg Pro Ala Ser Tyr Lys Lys Lys Ser

750

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									780	

Asp Ser Glu Val Lys Asn 780	
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Cys Phe Gly Glu Ser Glu Val Ser Trp Gln Tyr Pro Met Ser Glu Glu 50 55 60

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Phe Val Thr Val Leu Glu Val Ser Ser Ala Ser Ala Ala His Thr Gly 85 90 95

Leu Tyr Thr Cys Tyr Tyr Asn His Thr Gln Thr Glu Glu Asn Glu Leu 100 105 110

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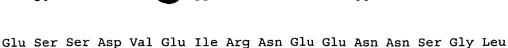
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- Phe Val Thr Val Leu Glu Val Ser Ser Ala Ser Ala Ala His Thr Gly
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- Leu His Asn Ser Glu Gly Val Val Pro Ala Ser Tyr Asp Ser Arg Gln
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- Gly Phe Asn Gly Thr Phe Thr Val Gly Pro Tyr Ile Cys Glu Ala Thr 180 185 190
- Val Lys Gly Lys Lys Phe Gln Thr Ile Pro Phe Asn Val Tyr Ala Leu 195 200 205
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- Tyr Lys Ser Gly Glu Thr Ile Val Val Thr Cys Ala Val Phe Asn Asn 225 230 235 240
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- Tyr Thr Leu Thr Val Pro Glu Ala Thr Val Lys Asp Ser Gly Asp Tyr
 275 280 285
- Glu Cys Ala Ala Arg Gln Ala Thr Arg Glu Val Lys Glu Met Lys Lys

Val Thr Ile Ser Val His Glu Lys Gly Phe Ile Glu Ile Lys Pro Thr Phe Ser Gln Leu Glu Ala Val Asn Leu His Glu Val Lys His Phe Val Val Glu Val Arg Ala Tyr Pro Pro Pro Arg Ile Ser Trp Leu Lys Asn Asn Leu Thr Leu Ile Glu Asn Leu Thr Glu Ile Thr Thr Asp Val Glu Lys Ile Gln Glu Ile Arg Tyr Arg Ser Lys Leu Lys Leu Ile Arg Ala Lys Glu Glu Asp Ser Gly His Tyr Thr Ile Val Ala Gln Asn Glu Asp Ala Val Lys Ser Tyr Thr Phe Glu Leu Leu Thr Gln Val Pro Ser Ser Ile Leu Asp Leu Val Asp Asp His His Gly Ser Thr Gly Gly Gln Thr Val Arg Cys Thr Ala Glu Gly Thr Pro Leu Pro Asp Ile Glu Trp Met Ile Cys Lys Asp Ile Lys Lys Cys Asn Asn Glu Thr Ser Trp Thr Ile Leu Ala Asn Asn Val Ser Asn Ile Ile Thr Glu Ile His Ser Arg Asp Arg Ser Thr Val Glu Gly Arg Val Thr Phe Ala Lys Val Glu Glu Thr Ile Ala Val Arg Cys Leu Ala Lys Asn Leu Leu Gly Ala Glu Asn Arg Glu Leu Lys Leu Val Ala Pro Thr Leu Arg Ser Glu Leu Thr Val Ala

Ala Ala Val Leu Val Leu Leu Val Ile Val Ile Ile Ser Leu Ile Val



Leu Val Val Ile Trp Lys Gln Lys Pro Arg Tyr Glu Ile Arg Trp Ile 545 550 555 560

Ser Pro Asp Gly His Glu Tyr Ile Tyr Val Asp Pro Met Gln Leu Pro 565 570 575

Tyr Asp Ser Arg Trp Glu Phe Pro Arg Asp Gly Leu Val Leu Gly Arg 580 585 590

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Gly Leu Ser Arg Ser Gln Pro Val Met Lys Val Ala Val Lys Met Leu 610 620

Lys Pro Thr Ala Arg Ser Ser Glu Lys Gln Ala Leu Met Ser Glu Leu 625 630 635 640

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Gln Arg Ser Leu Tyr Asp Arg Pro Ala Ser Tyr Lys Lys Ser Met 755 760 765

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Gly Leu Ala Arg Asp Ile Met His Asp Ser Asn Tyr Val Ser Lys Gly 835 840 845

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Trp Glu Ile Phe Ser Leu Gly Gly Thr Pro Tyr Pro Gly Met Met Val 885 890 895

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Pro Asp His Ala Thr Ser Glu Val Tyr Glu Ile Met Val Lys Cys Trp 915 920 925

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Val Glu Asn Leu Leu Pro Gly Gln Tyr Lys Lys Ser Tyr Glu Lys Ile 945 950 955 960

His Leu Asp Phe Leu Lys Ser Asp His Pro Ala Val Ala Arg Met Arg 965 970 975

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Ala Asp Ser Gly Tyr Ile Ile Pro Leu Pro Asp Ile Asp Pro Val

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Glu Ser Ser Asp Val Glu Ile Arg Asn Glu Glu Asn Asn Ser Gly Leu 65 70 75 80

Phe Val Thr Val Leu Glu Val Ser Ser Ala Ser Ala Ala His Thr Gly 85 90 95

Leu Tyr Thr Cys Tyr Tyr Asn His Thr Gln Thr Glu Glu Asn Glu Leu 100 105 110

Glu Gly Arg His Ile Tyr Ile Tyr Val Pro Asp Pro Asp Val Ala Phe 115 120 125

Val Pro Leu Gly Met Thr Asp Tyr Leu Val Ile Val Glu Asp Asp Asp 130 135 140

Ser Ala Ile Ile Pro Cys Arg Thr Thr Asp Pro Glu Thr Pro Val Thr 145 150 155 160

Leu His Asn Ser Glu Gly Val Val Pro Ala Ser Tyr Asp Ser Arg Gln 165 170 175

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Val Lys Gly Lys Lys Phe Gln Thr Ile Pro Phe Asn Val Tyr Ala Leu 195 200 205

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Val Ile Leu Ser Phe Glu Asn Asn Gly Asp Tyr Met Asp Met Lys Gln 725 730 735

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Lys Lys Ser Met Leu Asp Ser Glu Val Lys Asn Leu Leu Ser Asp 770 785

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Gln Val Ala Arg Gly Met Glu Phe Leu Ala Ser Lys Asn Cys Val His 805 810 815

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Pro Gly Met Met Val Asp Ser Thr Phe Tyr Asn Lys Ile Lys Ser Gly 900 905 910

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His Leu Ser Glu Ile Val Glu Asn Leu Leu Pro Gly Gln Tyr Lys Lys 945 950 955 960

Ser Tyr Glu Lys Ile His Leu Asp Phe Leu Lys Ser Asp His Pro Ala 965 970 975

Val Ala Arg Met Arg Val Asp Ser Asp Asn Ala Tyr Ile Gly Val Thr 980 985 990

Tyr Lys Asn Glu Glu Asp Lys Leu Lys Asp Trp Glu Gly Gly Leu Asp 995 1000 1005

Glu Gln Arg Leu Ser Ala Asp Ser Gly Tyr Ile Ile Pro Leu Pro 1010 1015 1020

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Ser Ser Ser Ser Thr Phe Ile Lys Arg Glu Asp Glu Thr Ile Glu 1055 1060 1065

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cag ctt tca tta ccc tct atc ctt cca aat gaa aat gaa aag gtt gtg Gln Leu Ser Leu Pro Ser Ile Leu Pro Asn Glu Asn Glu Lys Val Val 25 30 35
cag ctg aat tca tcc ttt tct ctg aga tgc ttt ggg gag agt gaa gtg Gln Leu Asn Ser Ser Phe Ser Leu Arg Cys Phe Gly Glu Ser Glu Val 40 45 50 55
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aga aat gaa gaa aac aac agc ggc ctt ttt gtg acg gtc ttg gaa gtg Arg Asn Glu Glu Asn Asn Ser Gly Leu Phe Val Thr Val Leu Glu Val 75 80 85
agc agt gcc tcg gcg gcc cac aca ggg ttg tac act tgc tat tac aac 703 Ser Ser Ala Ser Ala Ala His Thr Gly Leu Tyr Thr Cys Tyr Tyr Asn 90 95 100

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	_		 gag Glu	_	_	_		-					847
	_		 act Thr		_				_				895
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Tyr		Gly	gtg Val		Gly	Lys	Gly	Ile	Met				1183
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      The 'Xaa' at location 841 stands for Lys, Arg, Thr, or Ile.
<223>
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<222> (842)..(842)
<223> The 'Xaa' at location 842 stands for Lys, Asn, Arg, Ser, Thr,
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<223> The 'Xaa' at location 843 stands for Lys, Asn, Arg, Ser, Thr,
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<223> The 'Xaa' at location 846 stands for Lys, Asn, Arg, Ser, Thr,
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- <221> misc_feature
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- <223> The 'Xaa' at location 848 stands for Asn, Ser, Thr, Ile, Asp, Gly, Ala, Val, His, Arg, Pro, Leu, Tyr, Cys, or Phe.

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Cys Phe Gly Glu Ser Glu Val Ser Trp Gln Tyr Pro Met Ser Glu Glu 50 55 60

Glu Ser Ser Asp Val Glu Ile Arg Asn Glu Glu Asn Asn Ser Gly Leu 65 70 75 80

Phe Val Thr Val Leu Glu Val Ser Ser Ala Ser Ala Ala His Thr Gly 85 90 95

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Val Lys Gly Lys Lys Phe Gln Thr Ile Pro Phe Asn Val Tyr Ala Leu 195 200 205

Lys Ala Thr Ser Glu Leu Asp Leu Glu Met Glu Ala Leu Lys Thr Val



Tyr Lys Ser Gly Glu Thr Ile Val Val Thr Cys Ala Val Phe Asn Asn 225 230 235 240

Glu Val Val Asp Leu Gln Trp Thr Tyr Pro Gly Glu Val Lys Gly Lys 245 250 255

Gly Ile Thr Met Leu Glu Glu Ile Lys Val Pro Ser Ile Lys Leu Val 260 265 270

Tyr Thr Leu Thr Val Pro Glu Ala Thr Val Lys Asp Ser Gly Asp Tyr 275 280 285

Glu Cys Ala Ala Arg Gln Ala Thr Arg Glu Val Lys Glu Met Lys Lys 290 295 300

Val Thr Ile Ser Val His Glu Lys Gly Phe Ile Glu Ile Lys Pro Thr 305 310 315 320

Phe Ser Gln Leu Glu Ala Val Asn Leu His Glu Val Lys His Phe Val 325 330 335

Val Glu Val Arg Ala Tyr Pro Pro Pro Arg Ile Ser Trp Leu Lys Asn 340 345 350

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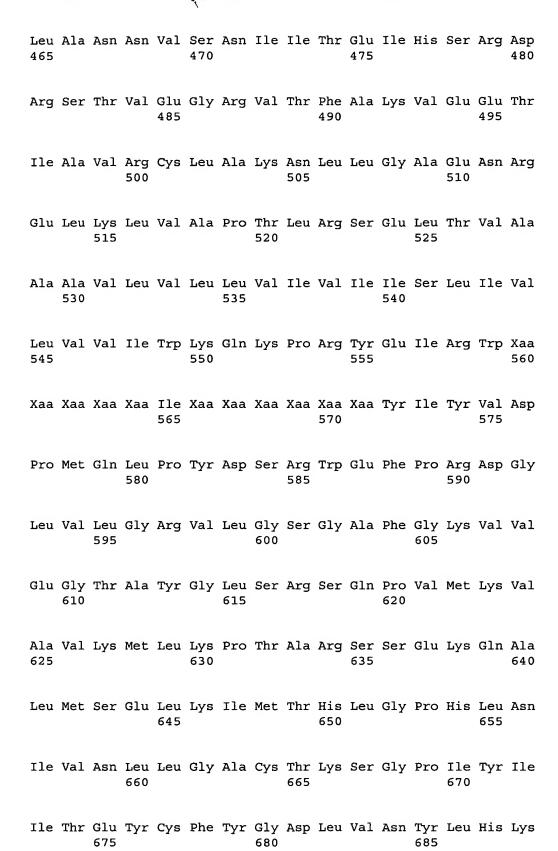
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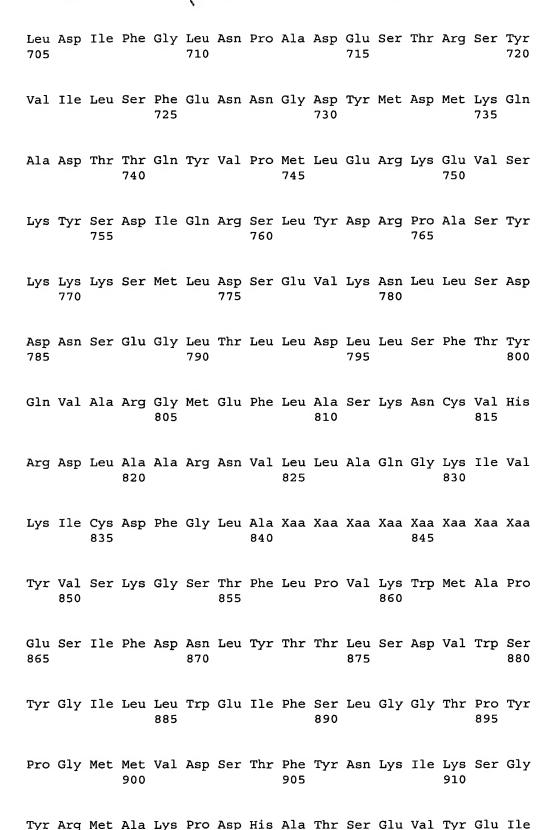
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Asn Arg Asp Ser Phe Leu Ser His His Pro Glu Lys Pro Lys Lys Glu



Met Val Lys Cys Trp Asn Ser Glu Pro Glu Lys Arg Pro Ser Phe Tyr



His Leu Ser Glu Ile Val Glu Asn Leu Leu Pro Gly Gln Tyr Lys Lys 945 950 955 960

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Glu Gln Arg Leu Ser Ala Asp Ser Gly Tyr Ile Ile Pro Leu Pro 1010 1015 1020

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Val Glu Asp Ser Phe Leu 1085

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